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Microbial Chitinases Purification: Conventional Protocols and Affinity Based Strategies (Review)

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Abstract: Microbial chitinases (EC.3.2.1.14) occupy a place of prominence among biocatalysts in nowadays biotechnology and have received tremendous interest among scientists and industrialists due to their wider range of biotechnological applications. Chitinases have been produced and purified to homogeneity from a large number of organisms ranging from bacteria and fungi to plant, animal and human; however the microbes still the major and the most industrially important source. In order to obtain chitinases in pure form for academic and biotechnological purposes, several purification protocols were proposed, this protocols varies in steps number, time and sequence, Furthermore they were laborious and cost and time consuming, for this reasons, The choice of the best purification protocol increasing the yields and folds still crucial and much recommended for enzyme characterization which help for understanding the biochemical aspects of microbial chitinases, in other hand the developing of an efficient large scale purification strategy remains a challenge to meet with increased industrial demands. The present review summarize the different purification protocols even conventional or affinity based, which have been employed for microbial chitinases and list the purification steps of each protocol comparing between them on basis of specific activity, yield and purification folds. As well as the recent available purification technologies also were reported. This review serves as an excellent literature reference and immense help for those working on microbial chitinases and planning for their purification.

Key words: Microbial Chitinase • Purification • Protocol • Chromatography • Affinity • Yield • Fold

INTRODUCTION

Chitinases (EC 3.2.1.14) are glycosyl hydrolases present in a wide range of prokaryotic and eukaryotic organisms, but industrially are produced by a number of microorganisms, including bacteria, actinomycetes, yeasts and fungi [1-4]. These enzymes are classified according to Henrissat and Bairoch [5] into subfamilies 18 and 19 on the basis of amino acid similarities within the catalytic domain.

Due to the expending applications of chitinases in various fields, it is important to understand their nature and properties for efficient and effective usage. In the last decades, vigorous research has been carried out on chitinases purification using different protocols varies in steps number, time and sequence, but the majority of them based on series of chromatography-based purification steps including ion exchange and size exclusion

chromatographies. Recently much attention has been focused on chitin affinity chromatography for chitinases purification [6, 7] due to their rapidity, simplicity and higher selectivity, capacity compared to the conventional protocols.

This review recapitulates the different protocols mostly cited in literatures concerning microbial chitinases purification using even conventional or chitin affinity based methods.

Chitinases Conventional Purification: A number of researchers have successfully purified and characterized Chitinases from a variety of microorganisms using different protocols. The crude extract subjected to precipitation using ammonium sulfate fractionation then followed by a series of chromatography-based purification steps including ion exchange, size exclusion chromatography to separate proteins on the basis of

Table 1: Chitinases Purification Using Conventional Protocols

Source / Reference	Purification steps	Yield (%)	Purification Fold
<i>Acinetobacter</i> sp. strain CHB 101 / [8]	Culture supernatant, (NH ₄) ₂ SO ₄ (0 – 80 %), CM-sepharose, Sephadex G-100	17 16	ND ND
<i>Aeromonas schubertii</i> / [9]	Cell filtrate, Ammonium sulphate 90 %, IEF	10	2
<i>Pseudomonas aeruginosa</i> K-187 / [10]	Culture supernatant, (NH ₄) ₂ SO ₄ (80%), DEAE Sepharose CL-6B, (NH ₄) ₂ SO ₄ (80%), Econo Pac q	27 7	10 6
<i>Cellulomonas flavigena</i> NTOUI / [11]	Culture filtrate, Q cartridge chromatography, hydrophobic interaction chromatography Superdex 75 HR	5.64	20.17
<i>Enterobacter</i> sp. NRG4 / [12]	Cell free supernatant, (NH ₄) ₂ SO ₄ (30 – 75 %), DEAE Sephadex, Sephadex G 200	31.1	44.12
<i>Serratia marcescens</i> BJL200 / [13]	Diluted periplasmic extract, phenyl-superose HR 5/5 (FPLC)	79	ND
<i>Vibrio alginolyticus</i> H-8 / [14]	Culture filtrate, crude chitinase, 2 nd DEAE-Toyopearl 650M, superdex 200 HR	10 6.7	8.3 14.5
<i>Xanthomonas</i> sp. strain Ak *ch B / [15]	Culture supernatant, (NH ₄) ₂ SO ₄ 100 %, phenyl Toyopearl 650 S, DEAE- Toyopearl 650M	NA 53	NA 16
<i>Bacillus circulans</i> WL-12 / [16]	Periplasmic fraction, (NH ₄) ₂ SO ₄ (40%), Q ceramic hyper D Ion exchange chromatography	90.6	33.7
<i>Bacillus stearothermophilus</i> CH ₄ / [17]	Culture filtrate, (NH ₄) ₂ SO ₄ (80%), DEAE cellulose, Butyl-Toyopearl, Sephadex 6-100, Mono-Q	1.31	387
<i>Streptomyces thermoviolaceus</i> OPC-520/ [18]	Culture filtrate, DEAE-Toyopearl 650 M, Sephadex G75, Phenyl Toyopearl 650 M, Mono-Q HR 5/5	29.9	20.6
<i>Streptomyces albobovineus</i> S-22 / [19]	Culture filtrate, Ammonium sulphate (80%), dialysis, Sephadex G200	40.07	2.3
<i>Metarhizium anisopliae</i> / [20]	Ammonium sulphate 85 %. DEAE Sephacel	2.74	2.5

ND: Not Determined

Table 2: chitinases purification using affinity based strategies

Source / Reference	Purification steps	Yield (%)	Purification Fold
<i>Aeromonas caviae</i> / [21]	Culture broth, H is Tag resin affinity. Cell extract, H is Tag resin affinity.	1.1 3.5	3 3
<i>Serratia marcescens</i> / [22]	Crude, colloidal chitin batch affinity, ultragel AC A54	ND	2 – 3
<i>Bacillus</i> sp. BG-11 / [23]	Cell free supernatant, (NH ₄) ₂ SO ₄ (50 – 80%), affinity binding to chitin, Sephadex G-100	15	16
<i>Bacillus</i> sp. strain MH-1 / [24]	Culture filtrate, chitin affinity (batch), chromate focusing	2.2 0.67 1.1	20 18 24
<i>Paenibacillus</i> sp. [25]	affinity adsorption on fermentation-processed chitin (FPC)	97	10.3
<i>Streptomyces venezuelae</i> P10[26]	80% Ammonium, sulfate precipitation, Affinity binding to chitin, DEAE-cellulose column	16.1	2.47
<i>Neurospora crassa</i> / [27]	- Crude, affinity precipitation with 0.5 % (w/v) chitosan solution - Crude, PEG-chitosan salt two phase system	85 86	27 34
<i>Rhizopus oligosporus</i> / [28]	Culture filtrate, Ammonium sulphate 90 %, chitin affinity column, Sephadex G-75, DEAE-Toyopearl	1.8 1.7	13.3 6

charge and size respectively. Comprehensive information with respect to purification steps used, yield and fold of purified Chitinases were briefly summarized in Table 1.

Chitinases Chitin Affinity Purification: Procedures that utilize the affinities of biomolecules and ligands for enzymes purification are gaining increasing acceptance due to their selectivity and higher capacity compared to the conventional protocols. Concerning chitinases, chitin affinity chromatography, has been shown to be a promising approach for chitinase purification. Chitinases from various microbial sources were purified to near

homogeneity using even batch or column affinity on chitin, colloidal chitin or chitosan with folds and yields reach to 34 and 86 % respectively [Table 2].

CONCLUSION

The choice of the best purification protocol increasing the yields and folds still crucial and chitin affinity chromatography is much recommended for chitinases purification, however extensive researches and further optimizations were needed for developing of an efficient large scale chitin affinity purification strategy to meet with increased industrial demands.

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